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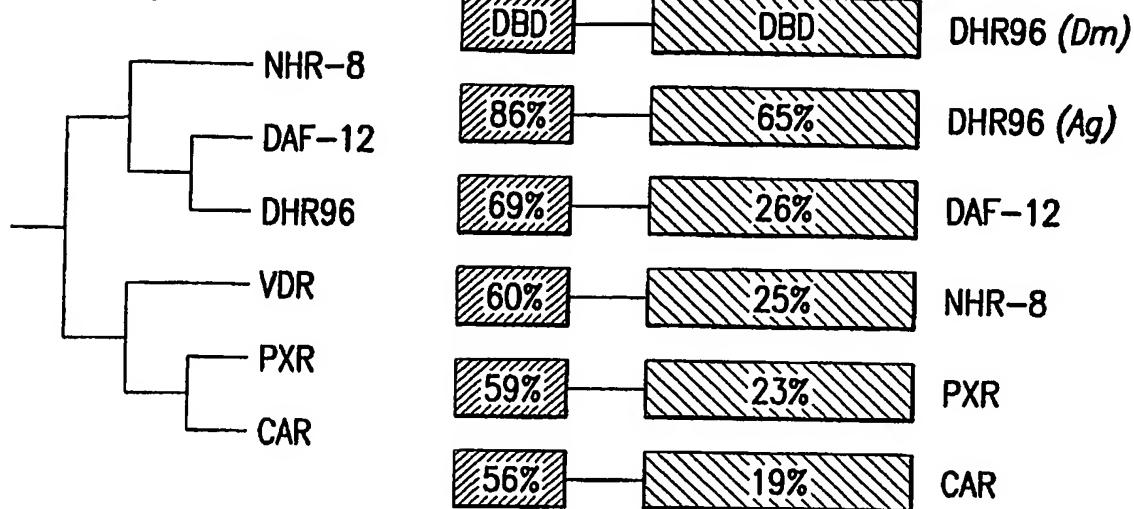


FIG.1A

FIG.1B

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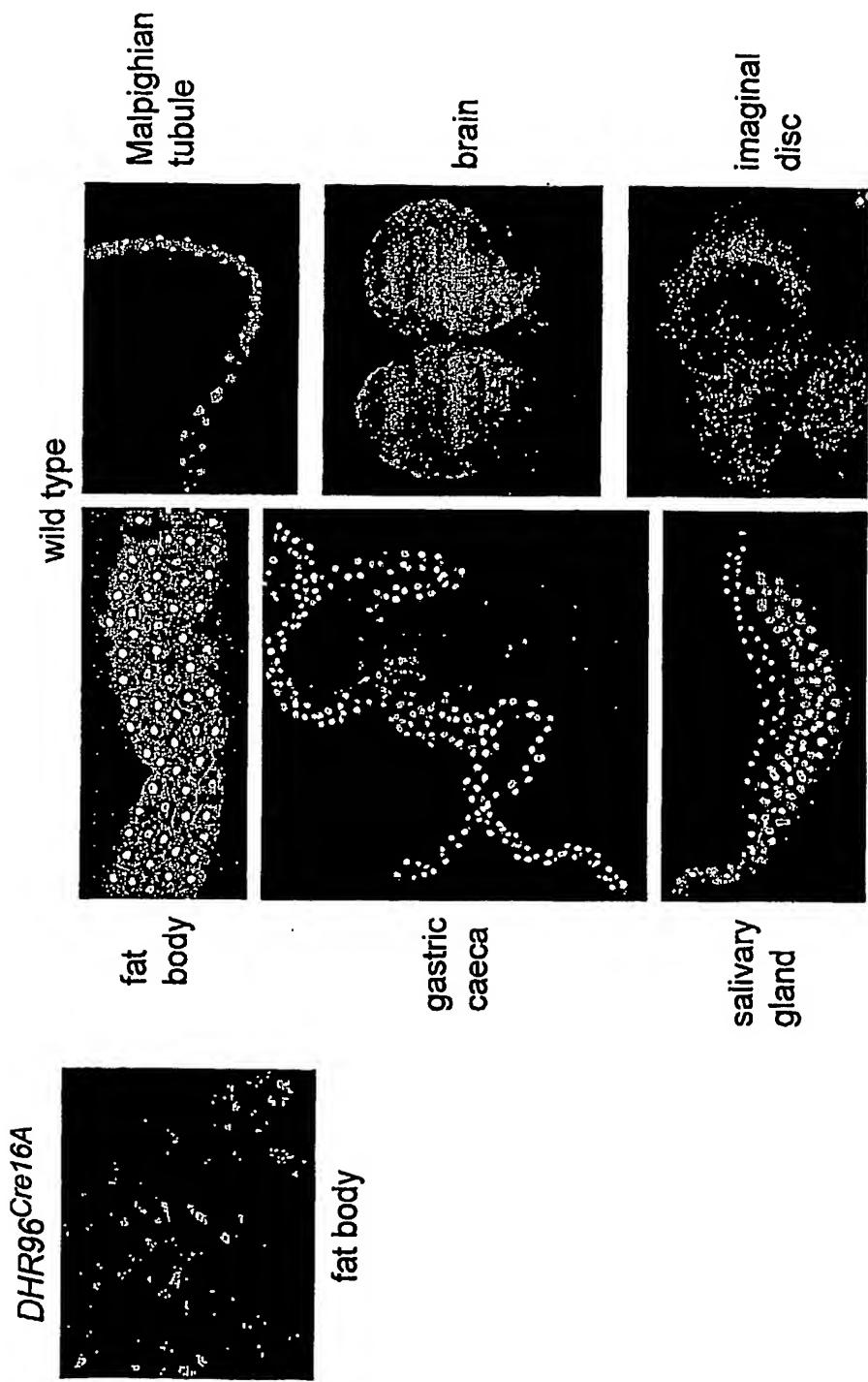


FIG.2

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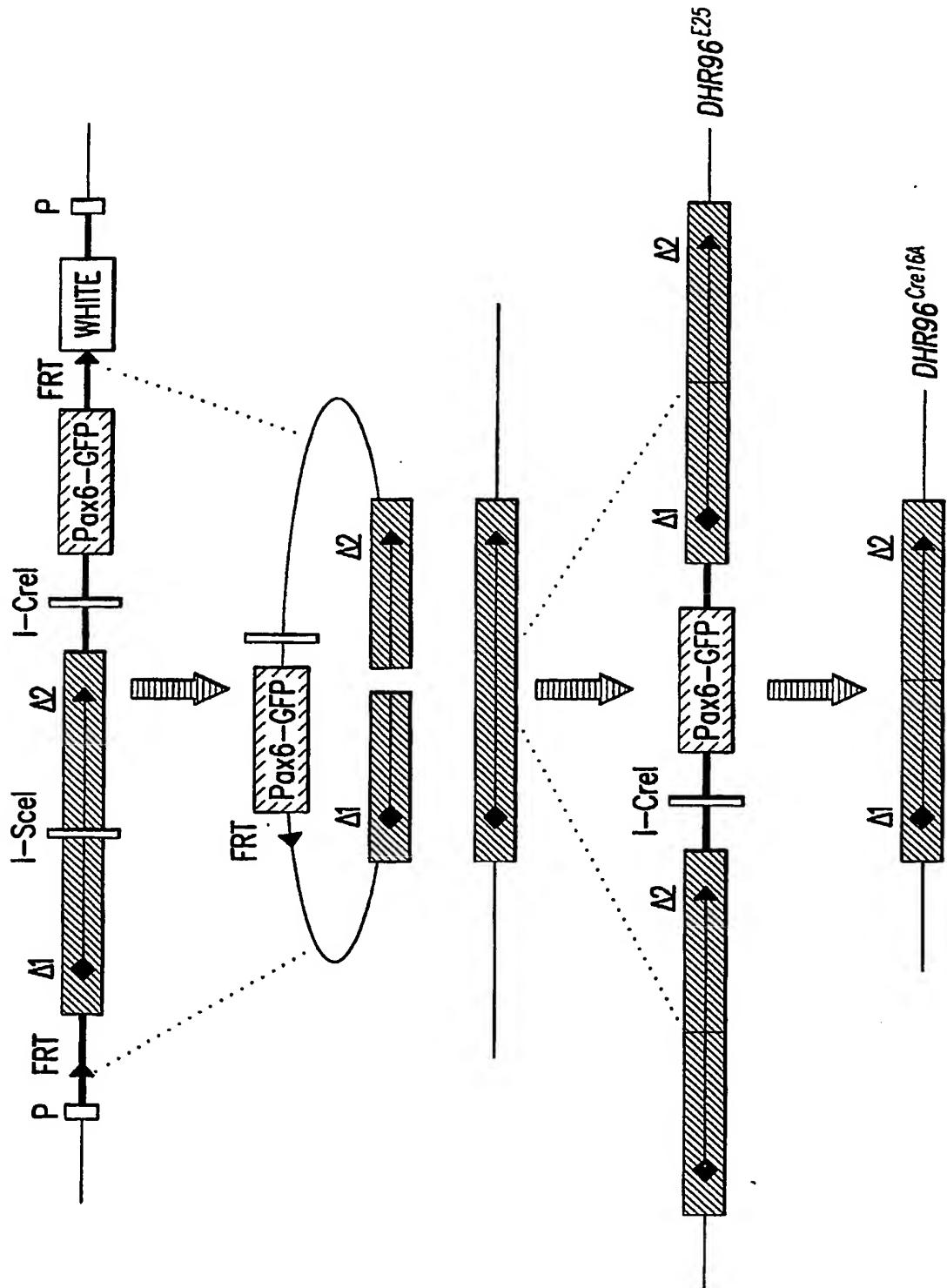


FIG. 3

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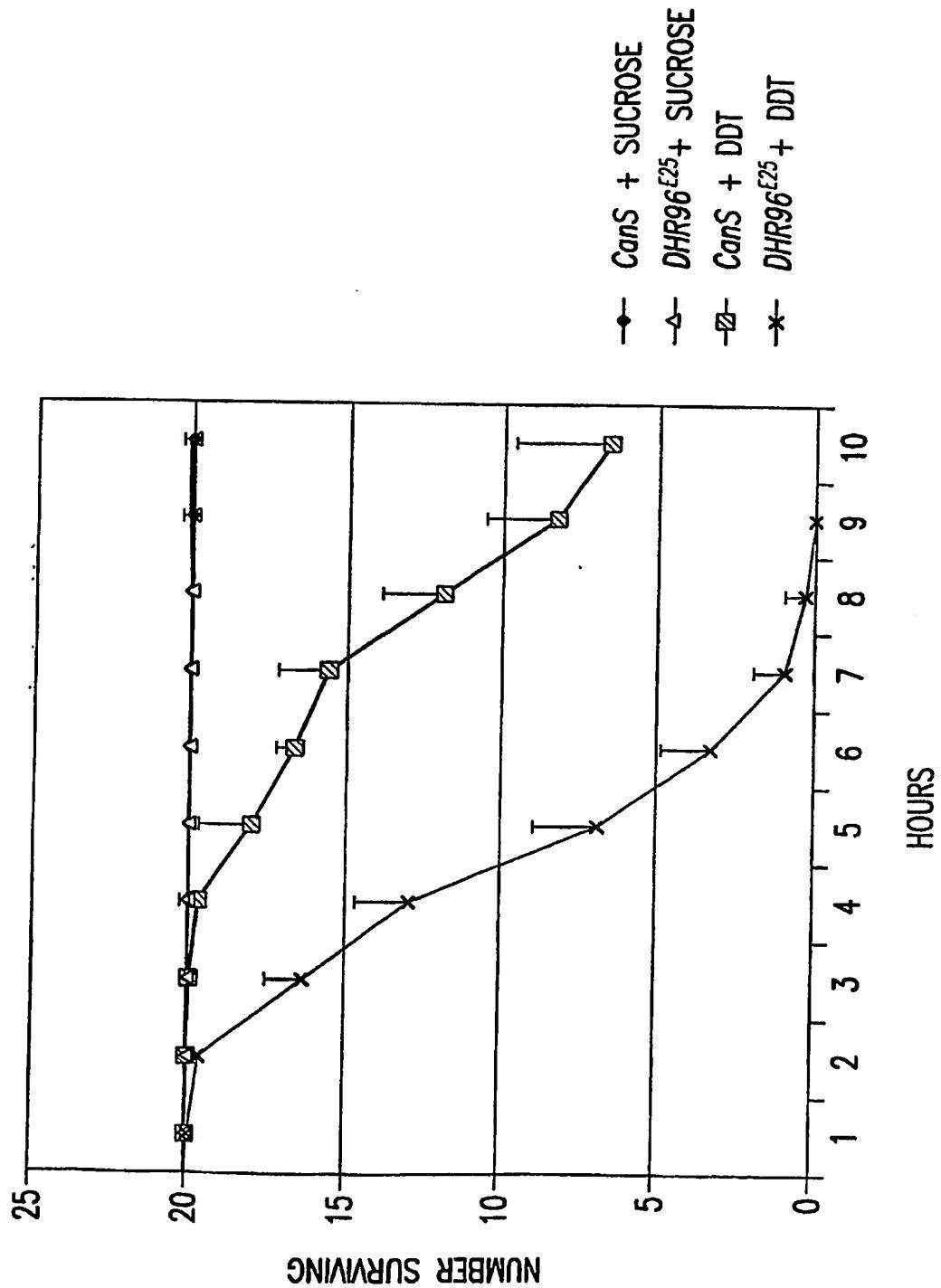


FIG. 4

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Usp	CSICGDRASGKHYGVYCEGCKGFFKRTVRKDLTYA-CRENR--NCIIDKRQRNRQCYCRYOKCLTGMKREAVQEERQ
SVP	CWCGDKSSGKHYGQFTCEGCKSFFKRSVRRNLTYS-CRGSR--NCPIDQHHRNQCQYCRLLKKCLKGMRREAVQGRV
dHNF4	CAICGDRATGKHYGASSCDGCKGFFRRSVRKNHQYT-CRFAR--NCWDKDKRNQCRCYCRLLKKCFKAGM
DHR3	CKVCGDKSSGVHYGVITCEGCKGFFRRSQQSSVNYQ-CPRNK--QCWDRVNRNRQCYCRLLKKCLKLM
DHR39	CPVCGDKVSGYHYGLLTCEGCKGFFKRTVQNKVYT-CVAER--SCHIDKTQRKRCPYCRFQKCLEVGM
E75	CRVCGDKASGFHYGVHCEGCKGFFRRSIQQKIQYRPCTKNQ--QCSILRINRNRCQYCRLLKKCIAVGM
E78	CKVCGDKASGYHYGVTCEGCKGFFRRSIQKQIEYR-CLRDG--KCLVIRLNRRNRCQYCRFKKCLSGM
EcR	CLVCGDRASGYHYNALTCEGCKGFFRRSVTKSAVYC-CKFGR--ACEMDMYMRRKCQECRLLKKCLAVGM
FTZ-F1	CPVCGDKVSGYHYGLLTCEGCKGFFKRTVQNKVYT-CVAER--SCHIDKTQRKRCPYCRFQKCLEVGM
T11	CKVCRDHSSGKHYGIYACDGCAGFFKRSIRRSLQYV-CKSQKQGLCVWDKTHRNQCRCRLLRKCFEVGM
DHR38	CAVCGDTAACOHYGVRTCEGCKGFFKRTVQKGSKV-CLADK--NCPVDKRRRNRCQFCRFQKCLWGM
DHR78	CLVCGDRASGRHYGAIISCEGCKGFFKRSIRKQLGYQ-CRGAM--NCEVTKHHRNRCQFCRLQKCLASGM
DHR96	CAVCGDKALGYNFNAVTCESCKAFFRRNALAKKQFT-CPFNQ--NCDITWTRRFQOKCRLRKOLDIGM
	unique region

FIG.5

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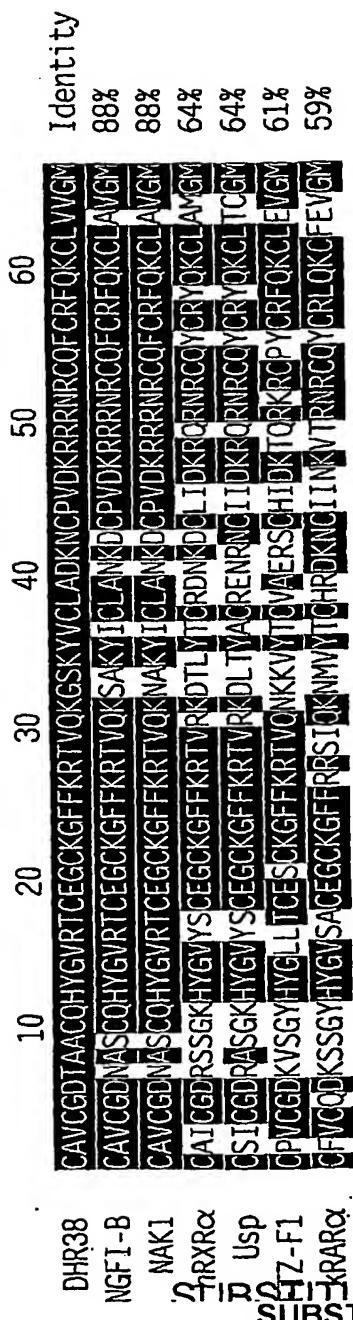


FIG. 6A

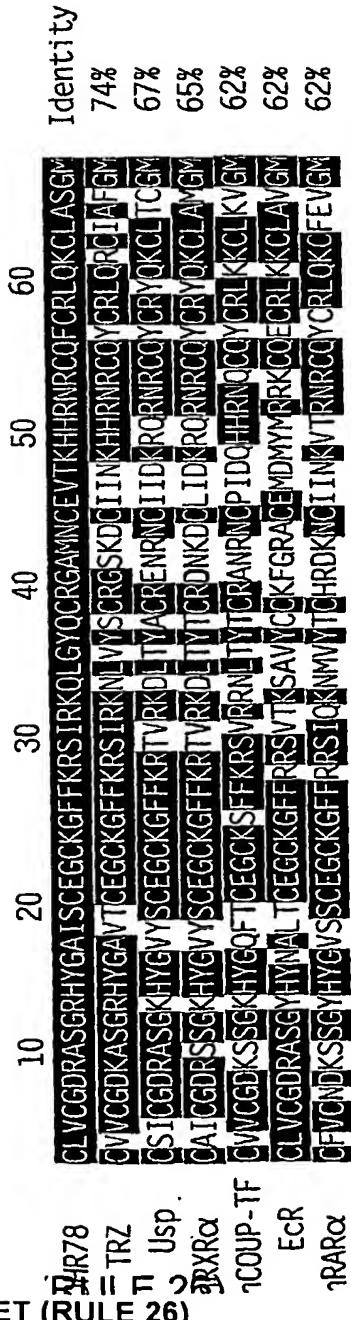


FIG. 6B

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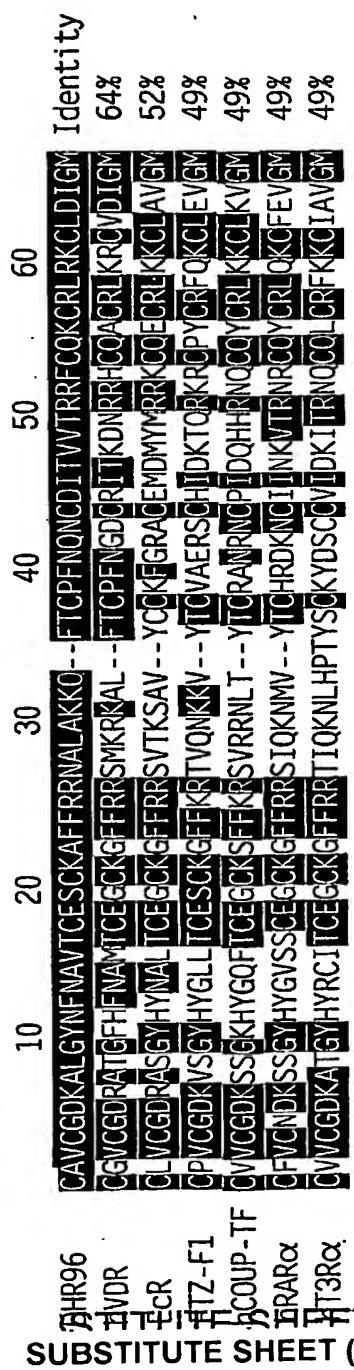


FIG.6C

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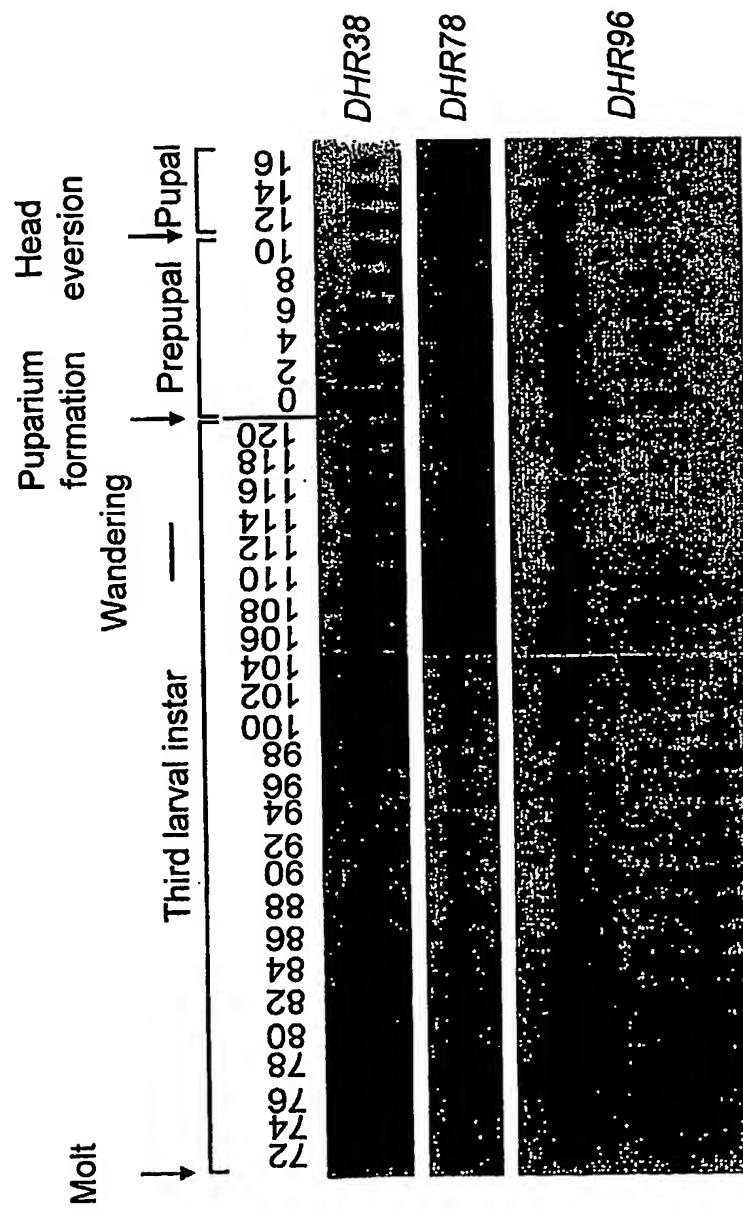


FIG. 7

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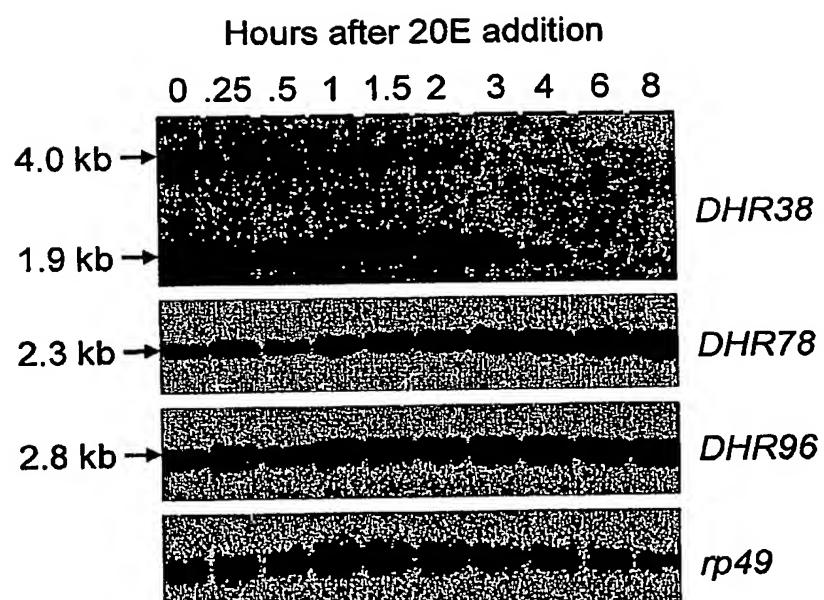


FIG.8

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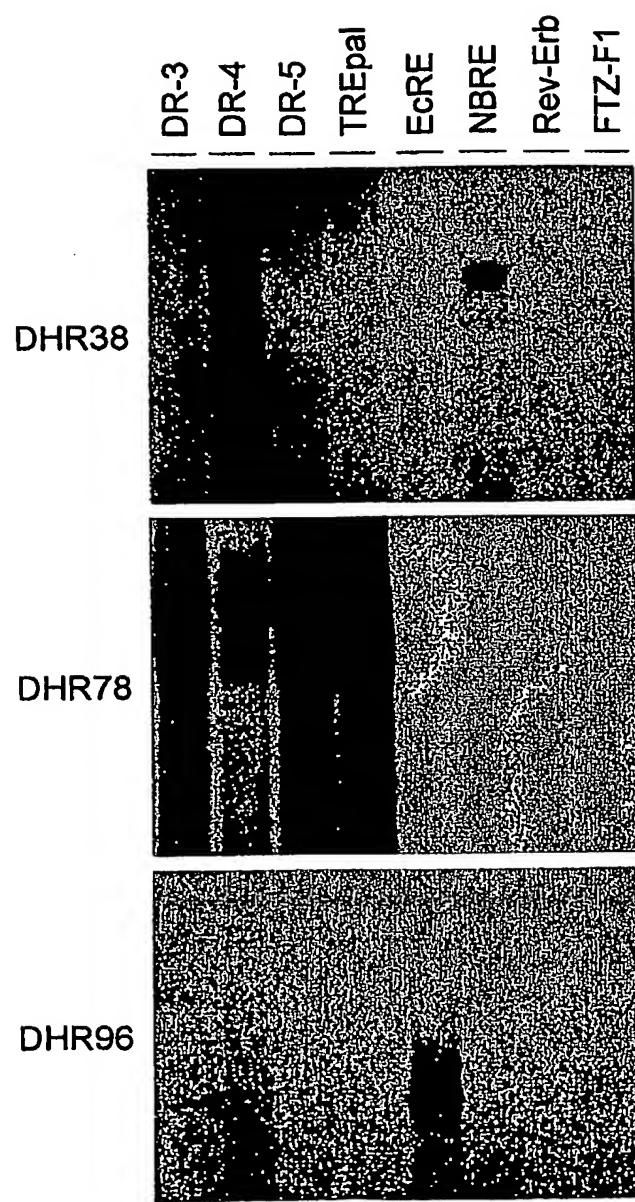


FIG.9

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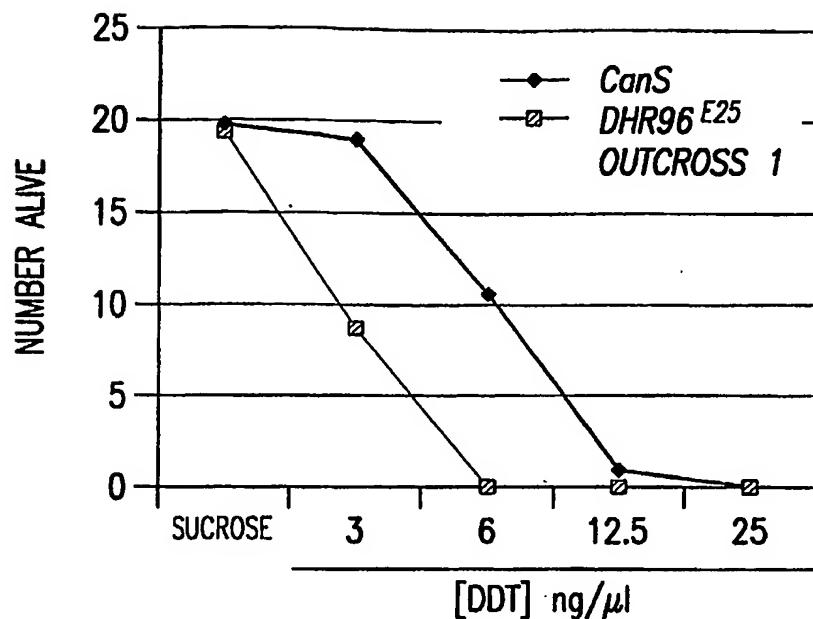
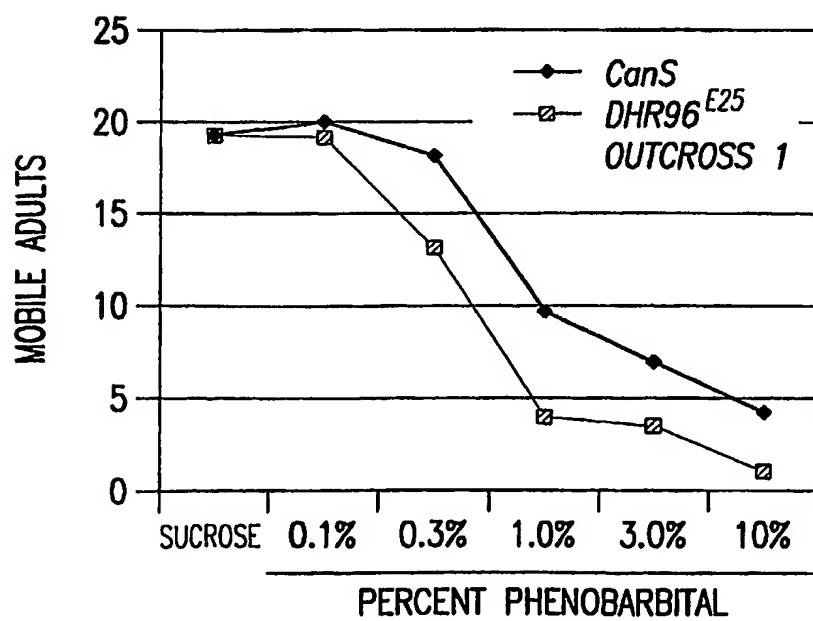


FIG. 10A

FIG. 10B  
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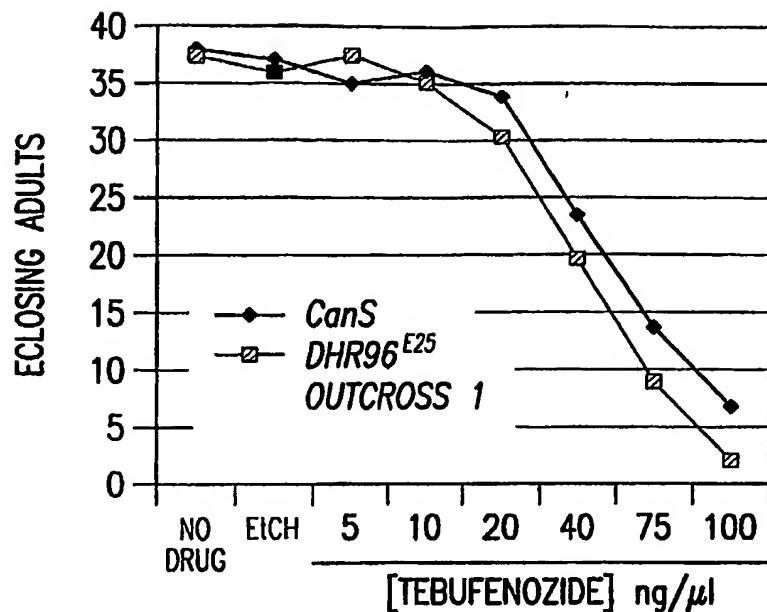


FIG.10C

scription		WPHS	96WPHS	fold change
3:FBgn0025644/sym=Lsp 1 gamma/prod=larval serum protein 1 gamma/prod=larval serum protein 1 gamma-s		11164.31	340.12	-32.82
3:FBgn0025311/sym=Lcp1/prod=larval cuticle protein1/prod=larval cuticle protein 1/func=structural protein c	3369.11	513.4	-6.56	
3:FBgn0019643/sym=Dat/name=Dopamine N acetyl transferase/prod= arylalkylamine N-acetyltransferase/func=	8868.78	1353.56	-6.55	
3:FBgn0033596/sym=CG7738/name=/prod=/func=/map=47E1-47E1/transc=CT23557/1er=405/GB:AE003:	4306.66	925.55	-4.65	
3:FBgn0030484/sym=CG16811/name=/prod=glutathione transferase-like/func=enzyme/map=11F1-11F1/transc=	946.93	221.04	-4.28	
2:FBgn0040609/sym=CG3348/name=/prod=/func=/map=97F6-97F6/transc=CT11231/1er=516/GB:AE003:	5850.94	1473.79	-3.97	
2:FBgn0039778/sym=CG18030/name=/prod=trypsin-like/func=endopeptidase/map=99F8-99F8/1transc=CT4:	1379.59	363.05	-3.8	
2:FBgn0034294/sym=CG5755/name=/prod=/func=/map=55B1-55B1/transc=CT18116/1er=1528/GB:AE00:	4602.49	1375.71	-3.35	
2:FBgn0039296/sym=CG10420/name=/prod=/func=/map=96C3-96C3/transc=T29244/1er=1794/GB:AE0	1271.55	405.9	-3.13	
2:FBgn0025333/sym=Lcp2/name=Larval cuticle protein 2/prod=larval cuticle protein 2/func=structural protein c	2050.81	723.13	-2.84	
2:FBgn0015039/sym=CG9522/name=Cytochrome P450, CYP623/func=cytochrome P45/map=51C	2526.49	947.68	-2.67	
2:FBgn0033398/sym=CG6223/name=/prod=cytochrome P450, CYP9B2/func=cytochrome P450	647.32	243.36	-2.66	
2:FBgn0033595/sym=CG18337/name=/prod=/func=/map=47E1-47E1/transc=CT41641/1er=540 /GB:AE00:	538.81	202.16	-2.66	
2:FBgn0036656/sym=CG_13026/name=/prod=/func=/map=73B5-73B5/transc=CT32244/1er=405/GB:AE00	2244.66	892.89	-2.51	
2:FBgn0000153/sym=b/black/prod=glutamate decarboxylase 2/func=glutamate decarboxylase : EC:4.1	7070.57	2889.86	-2.45	
2:FBgn0039798/sym=CG11313/name=/prod=nonphenol monooxygenase activator-like func=endopeptidase	1080.32	451.23	-2.39	
2:FBgn0032606/sym=CG17932/name=/prod=UDP-glucuronyl transferase/func=enzyme/map=36A9-36A9A	1214.19	515	-2.36	
2:FBgn0000594/sym=Est-P/name=Esterase P/prod=carboxyesterase/func=carboxyesterase: EC:3.1.1.1/map	475.26	205.79	-2.31	
2:FBgn0039239/sym=CG13641/name=/prod=/func=/map=98B4-96B4/1transc=CT33035/1er=429/GB:AE00:	1531.71	717.27	-2.14	
B:FBgn0014849/sym=Eig71Ei/name=/prod=/func=/map=71E4-71E4/1transc=CT22591/1er=3901GB:AE0035	1107.37	523.8	-2.11	
B:FBgn0034341/sym=CG17531/name=CG17531/prod=glutathione transferase/func=enzyme /map=55C9-55C9/1	3476.77	1645.19	-2.11	
B:FBgn0033830/sym=CG10814/name=/prod=/func=vitamin biosynthesis/map=50A6-50A6/1transc=CT30312	363.72	174.9	-2.08	

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FIG. 11A

FBgn ID	Symbol	Gene Name	Description	Product	Function	Map	Start	End	Length	Orientation	Chromosome	Link
3:FBgn0034010	sym=038157	/name=	serine protease 1	/prod=	/func=	/map=51F6-51F /transc=CT21959 /len=407	IGBA:ED031	1622.11	78.24	-2.06	2	
3:FBgn003356	sym=Ser990a	/name=	Serine protease 1/prod=serine endopeptidase	/func=	serine carboxypeptidase		15697.45	7724.85	-2.03			
3:FBgn001256	sym=Implt	/name=	Ecdysone-inducible gene L1	/prod=	ecdysone-inducible secreted membrane protein		1414.79	709.8	-1.99			
3:FBgn003046	sym=Pcp	/name=	upal cuticle protein/prod=	/func=	structural protein of pupal cuticle (Drosophila)		2111.66	1080.62	-1.95			
22FBgn003357	sym=Ser990b	/name=	Serine protease 2	/prod=	serine endopeptidase	/func=serine-type endopeptidase	15454.1	1955.78	-1.94			
5FBgn0036659	sym=039701	/name=	/prod=beta-glucosidase-1-like	/func=	ion channel	/map=73B5-73E5 /transc	1162.58	598.76	-1.94			
2FBgn0033821	sym=0G10799	/name=	/prod=	/func=	/map=50A1-50A1 /transc=CT30256 /len=591	IGB:AE00:	822.24	425.81	-1.93			
5FBgn0036024	sym=0G18180	/name=	/prod=	/func=	/map=67B10-67B10 /transc=CT41046 /len=887	IGB:AE	4368.44	2288.8	-1.91			
6FBgn0013307	sym=0dc	1	/name=Ornithine decarboxylase 1	/prod=	ornithine decarboxylase	/func=ornithine dcarboxylase	1007.84	530.1	-1.9			
6FBgn0029898	sym=0G14439	/name=	/prod=permease-1-like	/func=	transporter	/map=6C11-6C11 /transc=CT3	2393.73	1275.57	-1.88			
2FBgn0031653	sym=0G8871	/name=	/prod=	/func=	endopeptidase	/map=25B8-25B9 /transc=CT25448 /len=7	959.63	512.75	-1.87			
6FBgn0040565	sym=0G7606	/name=	/prod=	/func=	/map=9G05-90B5 /transc=CT23217 /len=321	IGB:AE003	8325.78	4504.75	-1.85			
6FBgn0031251	sym=0G4213	/name=	/prod=	/func=	motor	/map=21C2-21C2 /transc=CT13888 /len=3696	IGB:	146.23	19.37	-1.84		
8FBgn0033197	sym=0G17984	/name=	/prod=	/func=	/map=43E6-43E6 /transc=CT40154 /len=1417	IGB:AE0	1026.78	559.64	-1.83			
8FBgn0032229	sym=0G2229	/name=	/prod=serine protease-1-like	/func=	endopeptidase	/map=99F7-99F8 /trans	9394.5	5150.37	-1.82			
8FBgn0032889	sym=0G8331	/name=	/prod=glycerate dihydrogenase-1-like	/func=	enzyme	/map=38E9-38E9 /trans	3023.89	1676.12	-1.8			

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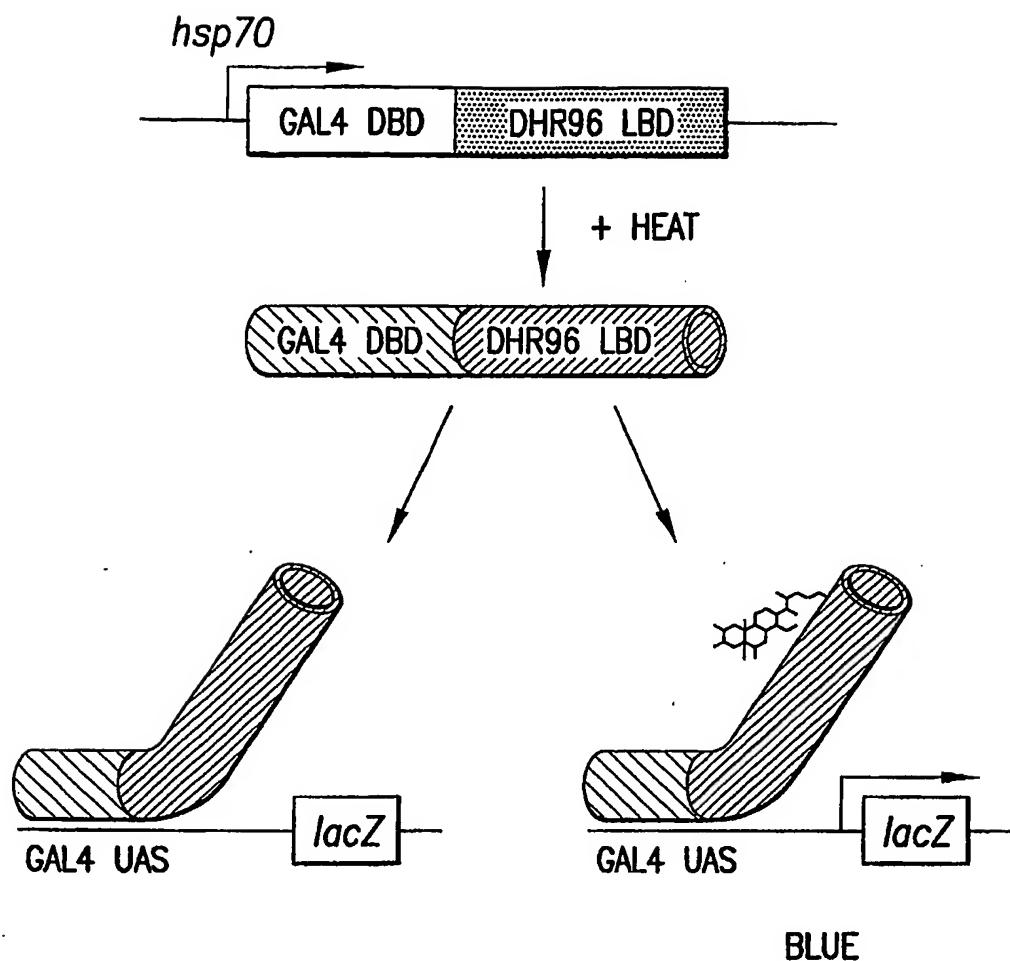


FIG. 12

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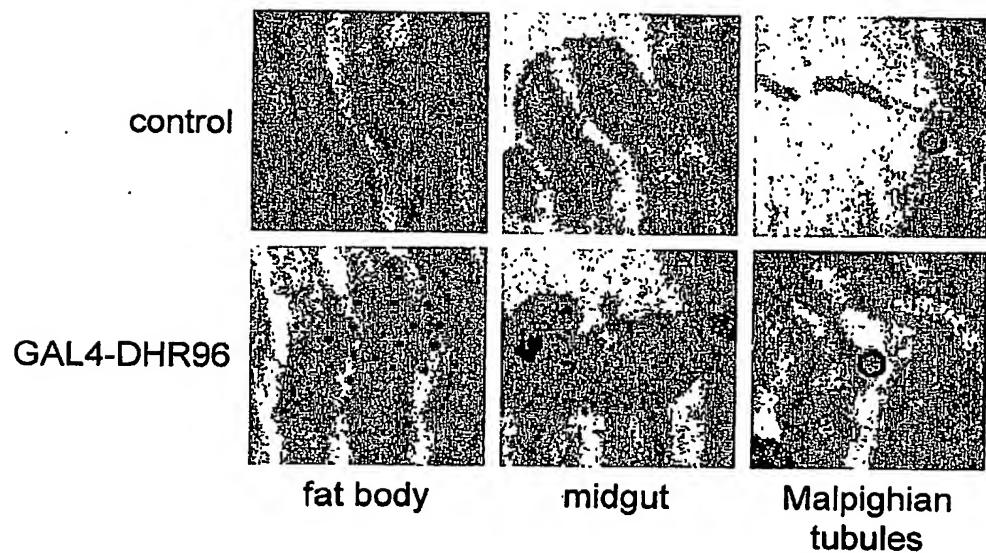


FIG.13